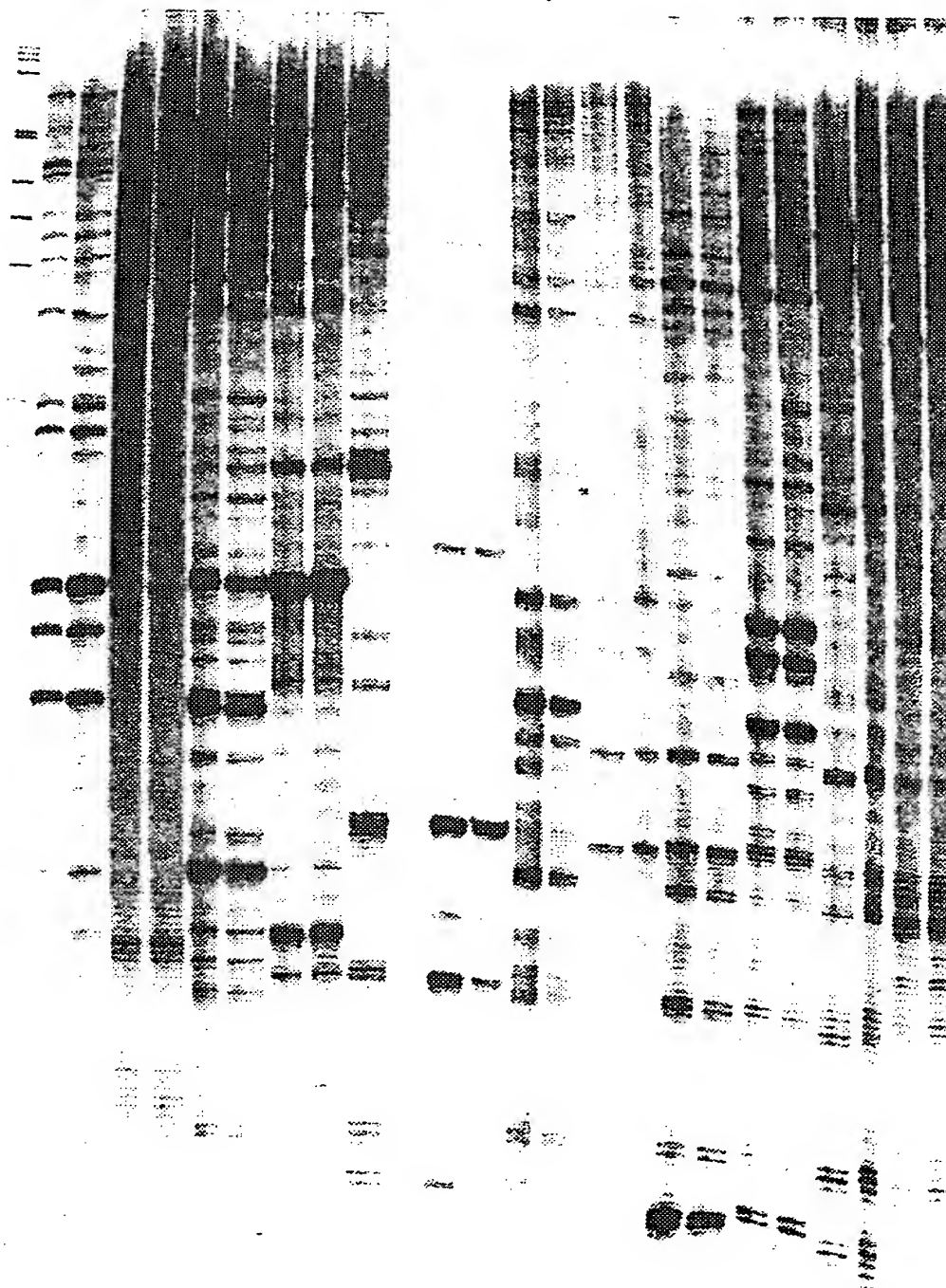
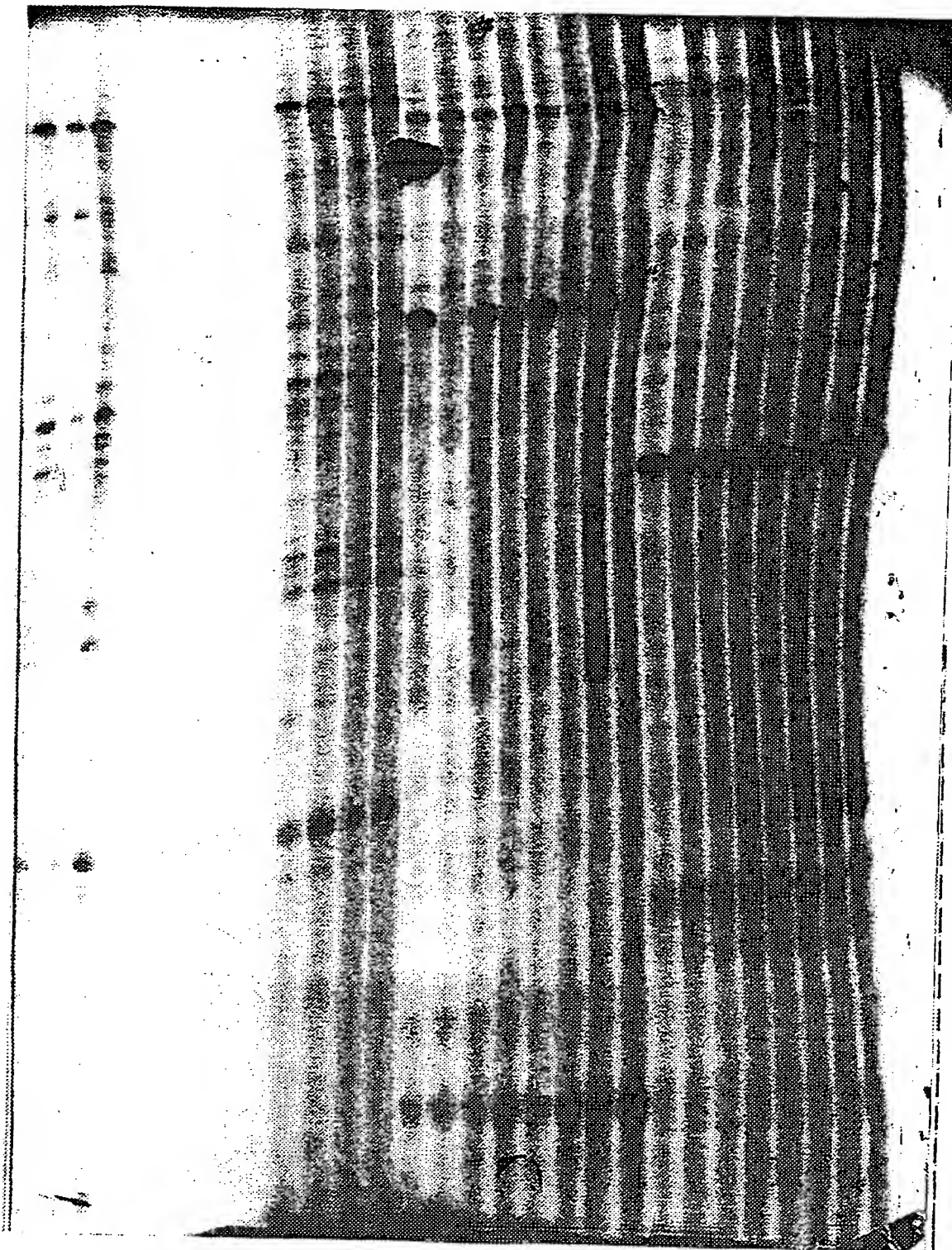


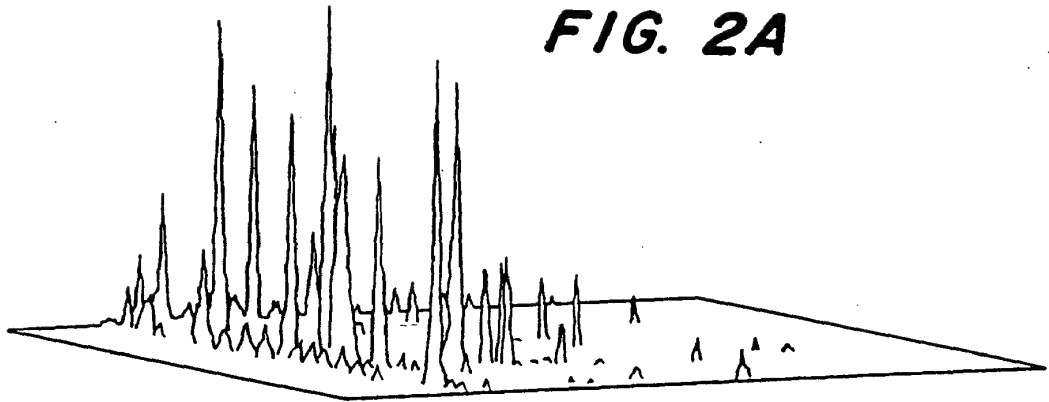
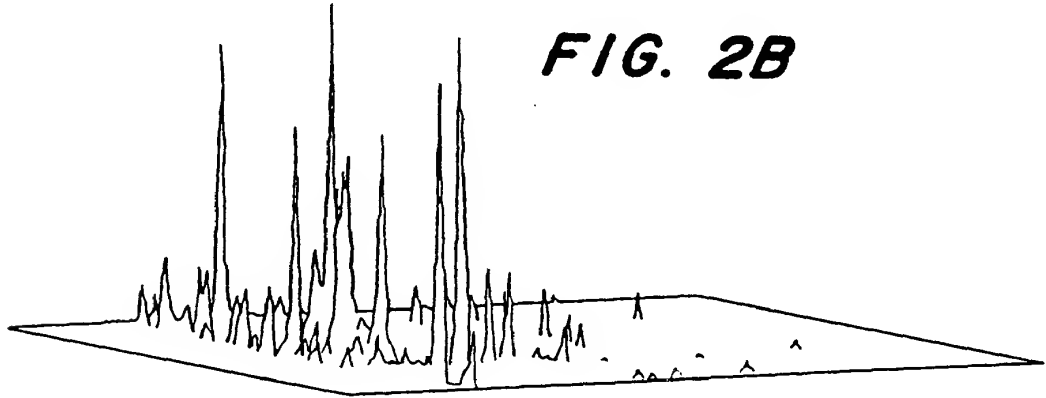
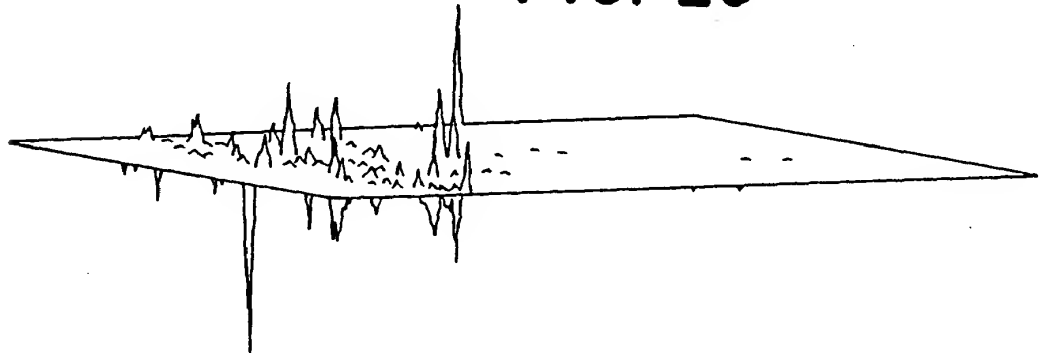
FIG. 1A



09674607.11401

FIG. 1B

09674607.11401

FIG. 2A**FIG. 2B****FIG. 2C**

* PARTIAL PATTERNS SHOWN REPRESENT APPROXIMATELY
5% OF COMPLETE PATTERN (ALL GENES EXPRESSED).

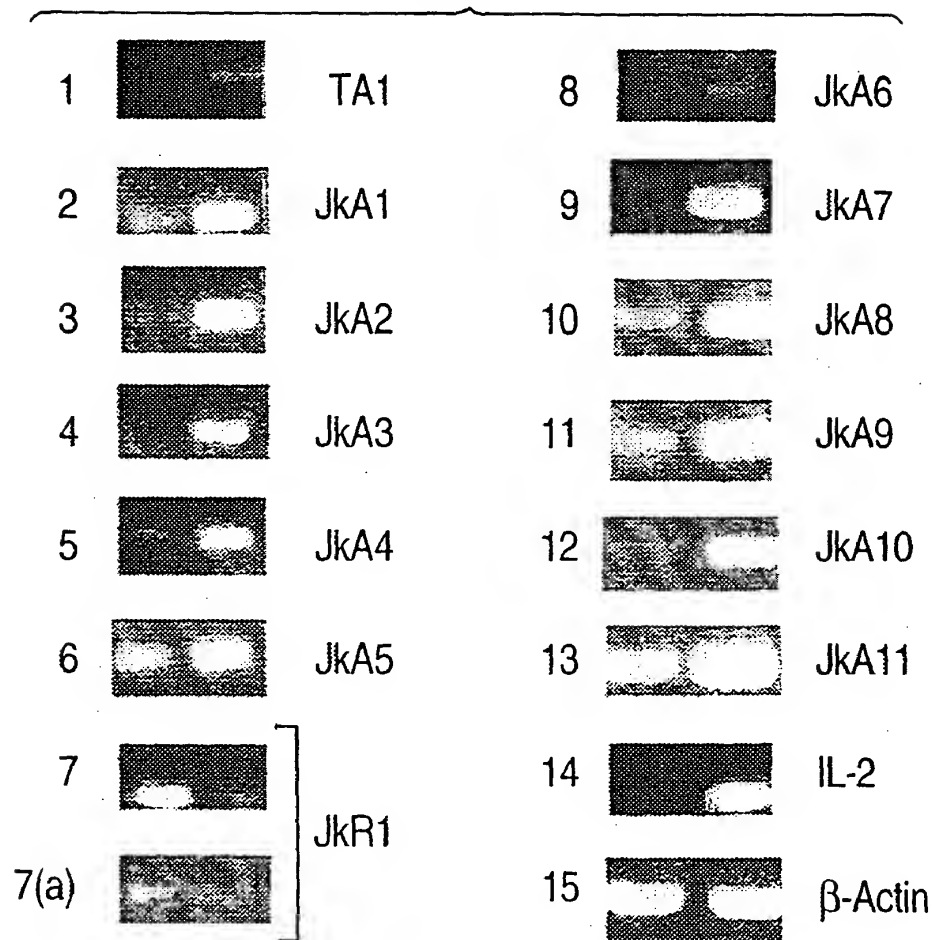
FIG. 3

FIG. 4

Differential Display Summary of Results					
No.	Clones	N1N2	Expression Patterns		Gene Bank Search and Analysis
			Control T cells	Activated Tcells	
1	TA1		0	÷	Unknown
2	JkA1		+	+++	Unknown
3	JkA2		+	+++	Unknown
4	JkA3		÷\-	++	Unknown
5	JkA4		÷\-	++	Unknown
6	JkA5		+	+++	Unknown
7	JkA6		0	+	Unknown
8	JkA7		0	+++	Unknown
9	JkA8		++	+++	Unknown
10	JkA9		++	+++	Unknown
11	JkA10		+1-	+++	Unknown
12	JkA11		++	+++	Human Serine Esterase
13			0	++	
14			0	++	Human Ribosomal Protein 58

FIG. 4 cont.

Differential Display Summary of Results				
No.	Clones	N1N2	Expression Patterns	
			Control T cells	Activated T cells
15			0	++
16			0	++
Gene Bank Search and Analysis				
Unknown				
Protein Translation Factor SVJ1				

FIG. 5

Clone v. Name	Enzyme	Band No.	R.P.	Level of Regulation*	Match (if any)	Percent ID	GenBank Accession	SEQ ID NO.
02_SV_011398_S AC7E_02	Bln I	11	10.2	4X I	Human mRNA for actin binding protein p57	98.182	HUMP57B	14
02_SV_011698_B SSIBAND26_02	Bln I	4	8.5	Absent	Human mRNA for KIAA0120 gene	99.194	HUMORFFA	15
04_SV102897_2A	Bln I	12	10.2	2X I	No human match			16
06_SV102897_1B	Bln I	5	8.5	1.5X I	Human ferritin light subunit mRNA	95.918	HUMFERLA	17
06_SV_011398_S AC14A_06	Bln I	13	10.2	2X I	Human STS WI-30177	98.611	G23064	18
08_SV102897_2B	Bln I	6	9.4	2X I				
10_NS108E_10	Bln I	14	10.2	1.5X I	Human DNA sequence from BAC 397C4	94.554	HS397C4	19
10_SV102897_1C	Bln I	7	9.4	Absent	Human clone 23732 mRNA	99.559	HSU79258	20
10_SV_011398_S AC15D_10	Bln I	15	10.2	3X I	Human hsc70 gene for 71 kd heat shock	98.785	HSWSC70	21
12_SV102897_2C	Bln I	18	10.3	2X I	Human colon carcinoma laminin-binding pr	99.602	HUMLAMB	
14_SV102897_1D	Bln I	9	9.4	2X I	No human match			22
16_SV102897_2D	Bln I	10	9.4	(Absent in R	Human mRNA export protein Rae1	98.788	HSU64720	23
18_SV102897_1E	Nsi I	8	8.4	2X I	No human match			24
19_SV011698_BS SIBAND15_19	Sac I	1	8.3	2X I	No human match			25

FIG. 5 cont.

Clone v. Name	Enzyme	Band No.	R.P.	Level of Regulation*	Match (If any)	Percent ID	GenBank Accession	SEQ ID NO.
20_SV102897_2E	Sac I	14	8.6	----	Human ferritin L chain mRNA	100	HUMFERL	26
23_SV011698_BS SIBAND17_23	Sac I	15	8.6	1.5X↓	Human mRNA for P1cdc47	100	HUMPICDC47	27
24_SV102897_2F	Sac I	7	8.4	Absent	Human pancreatic tumor-related protein	99.512	HUMPANCAN	28
26_SV102897_1G	BssSI	15	8.6	2X↓	Homo sapiens HLA-B gene	100	D83956	29
27_SV_011698_B SSIBAND19_27	BssSI	17	9.2	2X↓	Homo sapiens MLN50 mRNA	97.5	HSMLN50	30
30_SV102897_1H	BssSI	19	9.4	2X↓	No human match			31
31_SV_011698_B SSIBAND22_31	BssSI	22	10.2	4X↓	Human platelet activating factor acetyl/h	97.826	HSLIS10	32
37_SV_010898_S AC_BAND1_37	BssSI	26	10.3	2.5X↓	No human match			33

* Gene expression in Activated tissue is represented as (↑) or down (↓) regulated with respect to Resting Jurkat tissue. X represents fold increase or decrease in expression levels.